STIC-Biotech/ChemLib

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From:

Sent:

McGarry, Sean Tuesday, September 10, 2002 9:58 AM STIC-Biotech/ChemLib

To:

SEQ SEARCH Subject:

Sean McGarry 09/917,963 Office CM1 11D07 Mail box CM1 11E12 305-7028

Please, for, 09/917963, an oligomer search of SEQ ID NO:3. Please do NOT search EST data and do not search for interference. Thanks

> Point of Contact: **Toby Port** ecnnical Info. Specialist CM1 6A04 703-308-3534

Searcher: Phone:	
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TYPE OF S	EARCH:
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REFERENCE

AUTHORS

ACCESSION LOCUS HSMTP

RESULT

ALL EMBERIS

SOURCE KEYWORDS VERSION

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Score Match Length DB ID

Description

REFERENCE AUTHORS TITLE MEDITINE TITLE

Direct Submission

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/tissue_type="liver"
/clone_lib "Stratagene human liver cDNA"
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DRGKCKLQCSVEYPISSNVLVALLWPNPDGDDDQLLQTTMKDVNVENVNQQBCEKSIF
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/protein_id="CAA42200-1"
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1262 1200	1203 qaaqeeatttIqqaettIIIqqatttCaaaaqtqacaqeaqtatteteeteeaqqagagq 	<u> </u>
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	aareattttttetettttttägasttagasgreeasgsasaasgaaraassittlaa 3 	raogt teretaateaggaaaaaaaaaaaaaa aaaaaa aagaeereeaaara - 	t.t.cct.t.aaqcat.qacqqqa=3 	aaaaacaaaataaaaacaaaaceacacaaqqaqaacecaattitqtit. 9 	acagtcagaatagttattctc aagaggaaac	.pecaeaqegteattttqaateateatqtqaeqetttoaacaacqttctt.aqtt1a q 	.cuaaaaactgcagtttgatcaaatttgqqtatatgcagta_2 	ttqetetetqagaqeacaqeqtttacatatttacetqtattt 2 	tittaettaaattigteteeegadaggaaracaa 2 	agalqtqcaaagtqqtqtttqcccctcaqccqqaatagtacttccaqcqq=2 	attageaggatgtgauttreegeteratra 2 	COGGCAATTIGALHIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII	ctomataccoultettaqtitgcatecaratqacaa 2 	t ggestggaaacsagtacagaaasagaaacsaggettgga 2 	tgagtetaaaaeeegagtgaaaaatagagtgaetgtqgtaataaceaetgaeatcaeagt. 24 	tartteaggtgevaltgagtttagettgtggtgtetel 2 	######################################	HITTGATGTCCAAAATGCTGTCAGCATCTGGCGACCCTATCAGTGTGGTGAAAGGACTTAT 23
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                                                                                                                                                                                                                                                             205 REGIO AAAGTGGTTTGCCCCTCAGCCGGATAGTACTTCCAGCGGATGGTTTTGAAACTG 20597
                                                                                                                                                                                                                                                                                                                                                TRUL COLLANGGIAGANGGCACATAGAAATAACATC 3332
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20718 AGCTACAAAAAACTGCAGTFTGATCAAATTFTGGGFATATGCAGTATGCTACCCACAGCGT 20777
                                                                                                                                                                           ZUSYN AKURSTGATATITTAKITIGAATTIGTCICKKKAAAGGAKACAATGTGGCATGACTAAG 20657
                                                                                      2065) HACTIGUTCTCTGAGAGCACAGCGTTTACATATTTACCTGTATTTAAGATTTTTGTAAAA 20717
                                                                                                                                                                                                                                                                                     271/ atacagapteqtarinaceceteageequalagtacticcageqqaatqqttttqaaactg 2776
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30110
                     2897 agetaeaaaaactgeagtttgafeaaatttgggtatatgcagtatgetaeceacagegt 2956
                                                                                                       2777 unchytydatuttituchtyduttigictcccegadagggacaedatgiggeatgacidag 2836
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University, Institute of Genetics: 155 Li-Rong St. Section
Poiton, Taipei, Taiwan 11221, Republic of China
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                on Feb 6, 2001 this sequence version replaced gi:9929905
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (E-mail:ympetsalyym.edu.tw/ URL:http://genome.ym.edu.tw/
[e1:886-2-2826704], Fax:886-2-28264930)
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                                                                                                                                                                                                                                                                                                                   Collymore, A., Casarit, A., Trans, M., Collymore, A., Cooke, P., PeArcilano, K., Dewart, K., I. Collymore, A., Cooke, P., PeArcilano, K., Dewart, K., I. Dodde, S., Domino, M., Deyle, M., Perreira, P., Filth, Galdgan, J., Gardyna, S., Gindres, S., Goyette, M., Crah, Gardyna, S., Gindres, K., Beaford, A., B. Heaford, A., B. Heaford, C., Iliev, I., Bandardes, K., Jones, K., Macdonald, F., Klein, J., LaRocque, K., Lamadardes, K., Macdonald, F., Klein, J., LaRocque, K., Lamadardes, K., Macdonald, F., McCarthy, M., McEwan, P., McErik, A., McKernan, K., McMardy, T., Naylor, J., Mihova, I., Mitandarde, M., McMardy, T., Naylor, J., Mihova, I., Mitandarde, M., Pisani, C., Pollara, V., Kaymada, M., Kiley, K., Kostov Roy, A., Santos, B., Schauer, S., Severy, P., Spencer, Stange, Thomann, N., Stojansvier, N., Subramandah, A., Testaye, S., Theodore, J., Lirue, L., Liravers, M., Vassiliev, B., Zaimon, J., Zimmet, A., and Zhay, M., Xurarde, G., Zaimon, J., Zimmet, A., and Zhay, M., Xurarde, C., Zaimon, J., Zimmet, A., and Zhay, M.
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                                                                                                                                                                                                  Submitted (01-MAR-2000) Wintehead institute/MHT Consearch, 320 Charles Street, Cambridge, MA 02141 on May 6, 2000 this sequence version replaced ai:
                                                                                 http://itp.wensme.washib.n/ obj/PM/Poposimaekord
                                                                                                                                           All repeats were identified using RepealMasker: Smit, A.F.A. & Green, P. (1996-1997)
                                                                                                                                                                                                                                                                                                      Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (bases 1 to 175609)
Wob site: http://www-seq.wi.mit.edu
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Contact: sequence, submissioss genome.₩i.maf.e

Project Intoination

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* NOTE: This is a 'working draft' sequence. It currently consists of 22 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown.

* This record will be updated with the finished sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                               97500 110376: contig of 12877 bp in length 110377 110476: dap of 100 bp
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Sequencing worter MI3; M77815; 190% of reads
Chemistry: Dye-terminator Bid Dye; 100% of reads
Assembly program: Phrap; version 0 46074
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Center clone name 505_8_5
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43075 43174; gap of
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7679 12046: contig of 4868 bp in length
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4801 ____7578: contid of 2778 bp in length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (09-JAN-2012) Department of Genetics, Washington University, 4444 Forest Park Avenue, St. Louis, Missouri 63108, USA on Jan 9, 2002 this sequence version replaced 41.15922787.
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NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap between neighboring data submissions.

This sequence was tibished as follows unless otherwall regions were double stranded, sequenced with all chemistry, or covered by blue quality data (i.e., i.e., i.e., an attempt was made to resolve all sequencing as compressions and repeats: ill regions were coverage compressions. from more than one subclone: and the assembly was restriction digest

MAPPING INFORMATION

Mapping information for this state was provided by McPherson, Department of Genetics, Washington Univ sequence, see http://www. For additional information about the map posi-mence, see http://www.www.www.com/www.

SOURCE INFORMATION:

libraries. Genemics 51:1-8. The clone may be obtacesearch Genetics, Inc. (http://www.rosqen.com/) of and coworkers at the koswell back Cancer Institute approach for construction of pacterial artificial libraries. Genemics 51:1-m. The clone may be obdonor, as described by usocoawas.K., Woon, P.Y., Zha Tateno, M., Catanese, J.J. and to Jong, P.J. (1998) A The RPC1-11 human BAC librar; was made from the bi (http://bacpac.med.buffalo.edu)

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The clone sequenced to the celt is APOU1960; the the right is RP11-420012. Actual start of this coposition 1 of RP11-756F14; actual end is at base position NEIGHBORING SEQUENCE INFORMATIONS RP11-766F14.

Data from AC084421 was used to finish this close.

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                                                                                                                                                                                      Submitted (28-NeV 1994) D. Sharp, Bristol Myers Squibb, Hept of Metabolic Diseases, Princeton, New Jersey 08543-4000, USA
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Fax: 4157259689
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Metabolic Diseases, Princeton, New Jersey 08543-4000, USA
Location/Oualitiers
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Direct Submission
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Mammalia: Eutheria: Primates: Catarrhini: Hominidae: Homo.
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Direct Submission
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Sharp,D., Ricci,B., Kichele,P., Lib,M.C. and Worte
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sharp.D., Ricci.B., Kichzle,B., Lin,M.C. and Wetterau,J.R. Human microsomal trialyceride transfer protein large subunit gene
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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polypeptide for therapoutic use e.g. in reduction of obesity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     or cholesteryl ester in a mammalian species.
Diseases associated or affected by serum lipid levels, such as
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Local Similarity 99.8%;
Human; colon cancer; colon cancer antiqen; diagnosis; detection;
                          Human colon cancer antique encoding cDNA SEQ ID NO:725.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cancer-associated nucleic acid molecules (N) and proteins
the proteins are collectively known as colon cancer anti-
cancer antiques have cytostatic activity and can be used
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                                                                                                                                                                                                                                                                                                                                                                                               N.B. Pages 666 to 682 and page 70^{11} of the sequence first missing at time of publication, meaning no sequences are SEQ ID NO:1027 to 1052, 7921 and 7522.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    to express the proteins. N and P can be used in the prevand treatment of colorectal carcinomas and cancels. AABY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        diagnosis and treatment of diseases associated with image expression. For example, N and P may be used to treat dis
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                                                                                                                                    Sequence 1221 BP; 417 A; 214 A; 221 A; 464 T; 5 other:
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285 aacqttettaqtttaettafacefefefeada) eteaftiqqfacaqfeao
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mkNAs enceding secreted proteins. No ORF has yet been conclusively identified within the present sequence. The 5' ESTs were prepared from total human kNAs or polyA' kNAs derived from 30 different tissues. EST
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 225 BP; 82 A; 35 C; 32 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence is one of a large number of 5' ESTs derived from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-FEH-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21 FEB-2006; 2000EP 0200610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-SE 1 2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3167 ttuuqeatgaeyggaadaeeadaeeqtteeetauteaggaadaadaaaaaaaaaaa 3225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4
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                                                                                                                                                     l titaaguttittgtaaaaqctacaaaaactgcaqtttgatcaaatttqgqtatatgca 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              --
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                                                                                                                                                                                                                                                                                                               224: Conservative
                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                                           99.6%;
                                                                                                                                                                                                                                                                                                       Score 174; DB 21; Length 225; Pred. No. 2.4e-53; 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Giordano
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            r, 1 other,
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AAQ50446
                                                                                                                                                 Nucleic acid sequence encoding sub unit of MTP - used to express polypeptide for therapeutic use e.g. in reduction of obesity
                                                                                                                                                                                                                                                                                                                                                                                                 mutation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MTF; microsomal triglyceride transfer postein; obesity; atherosclerosis; detection; Inhibitor; prevention; stabilisation; reqression; lipid, serum etolesterol; so; triglyceride; 16; phosphatidyl choline; PC; cholesteryl ester; CE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ50446 standard; DNA; 302 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 72; 112pp; English.
                                                                                                                                          atherosclerosis etc.
                                                                                                                                                                                                                                                                      06-MAR-1992;
                                                                                                                                                                                                                                                                                            09 MAR 1993;
                                                                                                                                                                                                                                                                                                                   09-SEP-1993
                                                                                                                                                                                                                                                                                                                                         AU9334064-A
                                                                                                                                                                                                                                                                                                                                                                                                                                    exon
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MTP exon 13 mutant fragment.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAQ50446;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3059 aactagtidttigttaabaaacaaaaalaaaaacaadaaccacacacaaq 3103
                                                                                            Primers and probes were used in the detection of a gene detect
                                                                                                                                                                                      WPT: 1993-337245/43
                                                                                                                                                                                                                                  (WETT/) WETTERAU J
                                                                                                                                                                                                                                             (SQUI ) SQUIBB & SONS INC E R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         181 aactagtotttyttaaaaacaaaaataaaaacaaaaccacacacaaa 225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       J
                                                                                                                                                                                                            Sharp DY,
                                                                                                                                                                                                                                                                      9208-0847503.
                                                                                                                                                                                                                                                                                            93AU-0034064.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     285..302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                   119
                                                                                                                                                                                                                                                                                                                                                                                                            /number- 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "forward primer used to make probe"
                                                                                                                                                                                                                                                                                                                                                                 /note- "site of the C-to-T mutation"
                                                                                                                                                                                                                                                                                                                                                                             /label-
                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "reverse primer used to make probe"
                                                                                                                                                                                                                                                                                                                                                                                      /*tag=
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                                                                                                                                                                                                              Wetterau JR;
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                                                                                                                                                                                                                                                                                                                                                                             base 1830
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in an abetaliperreteinemic subject (intestinal biopsy). A C-to-T point mutation at base 1830 in exon 13 of the human MTP gene was revealed. This C to T mutation changes the norm

CGA arginine codon at residue 595 to a TGA translational stop

This C to T mutation changes the normal

signal, resulting in a 300 amino acid truncation of this protein. Bovine and human MTP are useful for the prevention, stabilisation or regression of atheroselerosis or for lowering the level of serum

lipids such as serum cholesterol, triglyceride, phosphatidyl choline

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110 AAQ 5.04.41
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  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           or cholestery) ester in a mammalian species.
Discusses associated or affected by serum lipid levels, such as
panereatitis, hyperdiyeemia, obesity, etc., may be treated.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 302 BP; 76 A; 67 C; 55 G; 104 F; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Miss microsomal triallyceride transfer protein; obesity; atherosclerosis; detection; inhibitor; prevention; stabilisat.on; regression; lipid; serum cholesterol; SC: triallyceride; TG; phosphotidy; choline; PC; confestery! ester: CE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAy50431 Standard; cDNA; 107 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bodine MIP 5' end.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10 MAY-1994 (tirst entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ50431:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   104 agrammatigic eqitegayit tetqaaqqaaatgqteyet cacaattatga eegitteetee 163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bos taurus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  164 unquiffgatettettetgeetacaetggetaeatagaaegta 206
                                                                                                                                                                                                                                                                                                                                                                                       misc_feature
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                                                                                                                                                                                                                                                                                                                                       AD9334064 A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity 100.0%;
                                                                                                                         Nucleic acid sequence encoding sub-unit of MTP - used to express
                                                                                                                                                       P PSDB; AAR42964
                                                                                                                                                                                           Gredd RE, Sharp DY, Wetterau JR;
                                                                                                                                                                                                                    (WEII/) WETTERAU J R
                                                                                                                                                                                                                                                            06 MAR-1992:
                                                                                                                                                                                                                                                                                                              09 SEP 1993.
                                                                                                              polypeptide for therapeutic use e.g. in reduction of obesity
                                                                                                                                                                                                                                 (SQUI ) SQUIBB & SONS INC E K
                                                                                                                                                                                                                                                                                     .5 MAR 1994;
The sequences given in \Delta \Delta Q 5.0429 \pm 3 may be used for the detection of MIP of MIP inhibitors.
                         Features Table)
                                    the knowledge of the 5' end of the bovine CDNA sequence (AAO50429) has been extended with the sequence given in AAO50431 (see
                                                                             rtaim 4-4; Page 31; 112pp; English
                                                                                                   atherosclerosis etc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ...
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                                                                                                                                                                   1993-337295/43.
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                                                                                                                                                                                                                                                                                                                                                                  /not e
                                                                                                                                                                                                                                                                                                                                                                                                                   /*t aq-
                                                                                                                                                                                                                                                                                                                                                                              pr.1*/
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100.0%; Pred. No. 9.2e-28;
                                                                                                                                                                                                                                                                                                                                                                "new sequence"
                                                                                                                                                                                                                                                                                                                                                                                                      "N-terminal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ); Mismatches
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Matches 76; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 107 BP; 30 A; 28 C; it 3; (8.1) 0 other:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 pancreatitis, hyperglycemia, chesity, etc., may be treate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                or cholesteryl ester in a manufalian species, piseases associated or affected by seriam lipid levels.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      adenosine triphosphate dependent kNA helicase; major histocompatibility complex class i histocompatibility pomplex class i histocompatibility phosphoglycerate kinase; immunoseppressive; immunosephosphoglycerate kinase; immunoseppressive; antidiatetic; an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human: single nucleotide polymorphism. SNP: andiopolethi
4-hydroxyburyrate: dehydroqunase: procein therapy:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAH79978;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-DEC-2000; 2000WO-US35346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-JUL-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                            Polymorphic nucleic acids encoding e.g. andiopoletin, a adenosine triphosphate-dependent PNA helicase and/or pakinase, useful for diagnosing and treating, e.g. cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI, 2001 418297/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-DEC-1999:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               990S-0472588
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100.0%; Pred. No.
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polymorphic variants of proteins (AA)98010-AA098240) reanglopoletin, 4-bydroxybutyrate, Johgdrowenaso, Jakenos (ATP)-dependent RNA helicase, major histocompatibility (ATP)-dependent BNIA helicase, major histocompatibility and igno and/or phosphnalyce class | histocompatibility and igno and/or phosphnalyce nucleic acid single nucleous ide polymorphisms (SNFs) an proteins have potential immunosyptocompative, immunostinous antirhoundatic, antisclerotic, antidiatetic, antibiliam antileukemic, neuroprotective and antibilerobial activity

The invention relates to nucleic acids (AAH79386-AAH80

1.11.10. 13.45 Claim 1; Page 233; 484pp: English.

diseases and infections

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%*83333333333333
Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diseases with a denetic component, such as autoimmune diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus crytheromatosus and Groote disease, inflammation, caucus (e.g. canners of the bladder, brain, breast, colon and kidney, lockemia), diseases of the nervous system, an intection of pathogenic organisms. They may also be used to alter phenotypic traits such as longerity, appearance,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 useful it dene/fritch therapy, racines, modulation of the expression and untivity of proteins related to angiopoletin. 4-hydroxybutyrate, dehydrouenase, adenosine triphosphate (AFP) dependent kNA helicase, major histocomparibility complex (MHC) Class I histocomparibility antiqen and/or phosphostyreate kindse. Disorders that may be prevented, diagnosed and/or treated by the above mothods include multificatorial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1816 ceattutteaagacatectacqtttqqaaatgcctqcaagcaaaattqtcc 1866
                                                                                                                                                                                                                                                                   09-SEP-1993
                                                                                                                                                                                                                                                                                                                                               polyA_site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MP: microsomal trialyzeride transfer protein; obesity; atheroselerosis; detection; inhibitor; prevention, stabilisation, retrossion: lipid; serum ecolesterol; 50, trialyzeride; TO, phosphatidyl choline; PC; cholesteryl ester; CB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ50429;
             atherosclerosis etc
                          polypeptide for therapeutic use e.g. in reduction of obesity
                                          Nucleic acid sequence encoding sub-unit of MTP - used to express
                                                                           P-PSDB; AAR42962.
                                                                                            WPI: 1993-337295/43
                                                                                                                       Gredd RE, Sharp DY,
                                                                                                                                                       (SQUI) SQUIBE & SONS INC E.E. (WEITZ) WEITERAU J.R.
                                                                                                                                                                                                        OH - MAR 1992;
                                                                                                                                                                                                                                    09-MAR-1993;
                                                                                                                                                                                                                                                                                                  AUS 434064-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bos taurus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    pancreatitis; hyperglycemia; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         strength, speed and endurance.
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    Mismatches

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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MTP; microsomal triglyceride transfer protein; obesity;
atherosclerosis; defection; inhibitor; prevention; stabilisation;
regression; lipid; serum cholesterol; SC; triglyceride; TG;
phosphatidyl choline; PC; cholesteryl estor; CE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             or cholesteryl ester in a mammalian species. Biscuscs associated or affected by serom lipid levels, such as pancreatitis, hyperalycemia, abesity, etc., may be treated.
                          The primers given in \Delta\Delta 0.504.34-4.2 and \Delta\Delta 0.504.44-4.5 are based on the normal buman cONA encoding the 88 kp component of MTP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ50443 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bovine and human MTP are useful for the prevention, stabilisation or regression of atheroselerosis or for lowering the lovel of serum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The sequences given in AAQS9429 33 may be used for the detection of MTP or MTP inhibitors.
                                                                        Disclosure; Page 68; 112pp; English.
                                                                                                      atherosclerosis etc.
                                                                                                                      polypeptide for therapeutic use e.g. in reduction of obesity
                                                                                                                                    Nucleic acid sequence encoding sub unit of MTP
                                                                                                                                                                  P-PSDB; AAR42965
                                                                                                                                                                                                              Gregg RE, Sharp DY,
                                                                                                                                                                                                                                          (SQUI ) SQUIBB & SONS INC E R. (WETT/) WETIERAU J R.
                                                                                                                                                                                                                                                                                        06-MAR-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              pascreatitis, hyperalycemia; abetalipoproteinemia; amplification
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in an abetalipoproteinemic subject (intestinal biopsy) The primers were used in the detection of a gene detect

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Boyline and human MIP are useful for the prevention, stabilisation or regression of afteroselerosis or for lowering the level of serum lipids such as serum choics erol, triddyceride, phosphatidyl choline of cholesteryl ester in a mammalian species.
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                                                                                                                                                                                                       number of human reproductive system related antiques. These can be used in the prevention and treatment of reproductive system disorders, including cancer. The present sequence is a genomic sequence encoding a
                            AAC25523 Standard; cDNA; 114 BP
                                                                                 2699 AGGANANANANANANANANANGTANGA 2671
                                                                                            Disclosure; SEC ID NO 7631; 1297pp + Sequence Listing; English
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17-N W-2000;
                                                                                                                                                                                                                                      The present investion provides the protein and coding sequences of
                                                                                                                                                                                                                                                                                 is used in preventing, treating or ameliorating a medical condition
                                                                                                                                                                                                                                                                                           Isolated nucleic acid molecule encoding a reproductive system antiqen
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                                                                                                                                                                                                                                                                                                                                     Barash SC,
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200008-0249217.
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                                                                                                                                             Length 9731;
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AAH71501
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3207 aaaaaaaaaaaaaaaaaaaagtaagac 3233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          derived from the 5' ends of mkHAs and even in those cases where longer cDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mkNAs with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and for diagnostic, forensic, gene therapy and chromosome mapping procedures
W0200142467-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID 29598; 71pp + CD:ROM; English
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                                                                                                                                                  Cervical cancer; cytostatic; pre-malignant condition; gene therapy; ss
                                                                                                                                                                                                                                 Human cervical cancer marker nucleic acid 2775.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAH71501 standard; cDMA; 203
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                                                                                                                                                                                                                                                                                                            19 SEP 2001
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100.0%; Pred. No. 2.8;
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county Match
Rest Local Similarity 100.0%;
Matches 27: Conservative (
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21-DEC-1999;
14-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                useful for dene therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WP1: 2001-475006/39
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                                                                                                                                                                                                                                                                                                                                                                                                              AAH70076 Standard; cbNA; 225 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 203 BP; 91 A; 25 C; 29 G; 58 T; 0 other;
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                                                                                                                                                                                                                                                                                           Pervical cancer: cytostatic; pre-maliquant condition; gene therapy;
                                                                                                                                                                                                                                                                                                                       Human cervical cancer marker nucleic acid 1350
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             Schlegel K. Dreds J,
                                                                                                    12 MAY 2000;
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                                         (MILL:) MILLENNIUM PREDICTIVE MEDICINE INC
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200000S-0203791
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9908-0171350
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               Berger A.
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Pred. No.
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RESULT 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cervical cancer with cytostatic activity. The nucleic acto
polypeptides are useful: to assess it a patient is attrict
cervical cancer or has a pro-maillanant condition; to monif
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WP1; 2001-375006/39
                                                               human bladder tumour and which have symmetatic activity for recombinant expression of (1) and to isolate complete used to identify agents suitable for treatment of bladder directly treat this form of cancer (including expression).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 225 BP: 92 A: 33 C: 37 G: 1. 1. 1. Other
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                                                                                                                                                                                                                                          New nucleic acid sequences expressed in bludder tumor dis-
derived polypeptides, for treatment of cladder tumor and
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                                                                                                                                     This invention describes novel polypoptide frauments (1) polynuclectides (II) that encode them that are highly exp
                                                                                                                                                                                         Claim 3; Page 90; 132pp; German.
                                                                                                                                                                                                                             of therapeutic agents -
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treatment; dene therapy; ESI; ss.
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hes 27: Conservative t
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Pred. No. 2.5;
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therapy vectors) or are used in a preparation for cancer is also used for the generation of specific antibodics. (identified by assembling PSTs (expressed sequence task) I particular tissue type before comparison of expression par

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Best Local Similarity
Matches 27; Conserv
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24 - FEB - 2009;
24 - FEB - 2000;
C2 - MAR - 2000;
16 - MAR - 2000;
17 - MAR - 2000;
19 - MAY - 2000;
19 - MAY - 2000;
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       Homo supieus.
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cancers and metastases Nucleic acids encoding 3224 haman nervous system antigen polypeotides, useful for preventing, digmussing and/or treating nervous system

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(b) immune disorders e.q. Addison's disease, allergies, autoimmune haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, multiple selergis, rheumateid artheritis and electative chilis: (c) cardiovascular disorders such as mycardial ischaemias; (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and chilepsy: and (l) intections diseases such as viral, bacterial, fungal and parasitic intections. the invention relates to movel genes (ARAll)04-ARA21534) and proteins (ARAll)4578 ARRIB001) useful for preventing, treating or ameliorating modical conditions e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant)agonists are useful in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer and other cancers of the adrend gland, bone, bone marrow, breast, gastrointestinal fract, liver, lung, or urogenical; Note: The sequence data for this parent did not form part of the printed specification, but was obtained in electronic format directly from when at Hp.wipo.int/pub/published_pet_sequences.

gordy Match Best Local Similarity 100.0%; 0.8%; Score 27; DB 22; Pred. No. 1.8; Length 1594;

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